

sequence comparison
A

10608449Search.txt

RESULT 2
AX086664
LOCUS AX086664 2406 bp DNA linear PAT 09-MAR-2001
DEFINITION Sequence 616 from Patent WO0112659.
ACCESSION AX086664
VERSION AX086664.1 GI:13276007
KEYWORDS
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1
AUTHORS Wiemann, S.
TITLE Human dna sequences
JOURNAL Patent: WO 0112659-A 616 22-FEB-2001;
German Human Genome Project (DE)
FEATURES
source Location/Qualifiers
1. 2406
/organism="Homo sapiens"
/mol_type="unassigned DNA"
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ORIGIN

Alignment Scores:

Pred. No.:	2.64e-244	Length:	2406
Score:	3176.00	Matches:	595
Percent Similarity:	100.00%	Conservative:	0
Best Local Similarity:	100.00%	Mismatches:	0
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DB:	6	Gaps:	0

US-10-608-449-4 (1-595) x AX086664 (1-2406)

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Qy	21	ProPheProSerIleLysAsnGluSerAsnTyrHisProPhePhePheArgThrArgAla	40
Db	406	CCTTTTCCTTCCATTAAAAACGAAAGCAATTACCACCCTTCTTCTTTAGAACCCGAGCC	465
Qy	41	CysAspLeuLeuLeuGlnProAspAsnLeuAlaCysLysProPheTrpLysProArgAsn	60
Db	466	TGTGACCTGTTGTTACAGCCGGACAATCTAGCTTGTAACCCCTTCTGGAAGCCTCGGAAC	525
Qy	61	LeuAsnIleSerGlnHisGlySerAspMetGlnValSerPheAspHisAlaProHisAsn	80
Db	526	CTGAACATCAGCCAGCATGGCTCGGACATGCAGGTGTCCTTCGACCACGCACCGCACAAAC	585
Qy	81	PheGlyPheArgPhePheTyrLeuHisTyrLysLeuLysHisGluGlyProPheLysArg	100
Db	586	TTCGGCTTCCGTTCCTTCTATCTTCACTACAAGCTCAAGCACGAAGGACCTTCAAGCGA	645
Qy	101	LysThrCysLysGlnGluGlnThrThrGluMetThrSerCysLeuLeuGlnAsnValSer	120
Db	646	AAGACCTGTAAGCAGGAGCAAACTACAGAGATGACCAGCTGCCTCCTTCAAAATGTTTCT	705
Qy	121	ProGlyAspTyrIleIleGluLeuValAspAspThrAsnThrThrArgLysValMethis	140
Db	706	CCAGGGGATTATATAATTGAGCTGGTGGATGACACTAACACAACAAGAAAAGTGATGCAT	765
Qy	141	TyrAlaLeuLysProValHisSerProTrpAlaGlyProIleArgAlaValAlaIleThr	160

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Db	766	TATGCCTTAAAGCCAGTGCCTCCCCGTGGGCCGGGCCCATCAGAGCCGTGGCCATCACA	825
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Db	826	GTGCCACTGGTAGTCATATCGGCATTGCGACGCTCTTCACTGTGATGTGCCGCAAGAAG	885
Qy	181	GlnGlnGluAsnIleTyrSerHisLeuAspGluGluSerSerGluSerSerThrTyrThr	200
Db	886	CAACAAGAAAAATATATATTACATTTAGATGAAGAGAGCTCTGAGTCTTCCACATACACT	945
Qy	201	AlaAlaLeuProArgGluArgLeuArgProArgProLysValPheLeuCysTyrSerSer	220
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Qy	221	LysAspGlyGlnAsnHisMetAsnValValGlnCysPheAlaTyrPheLeuGlnAspPhe	240
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Db	1066	TGTGGCTGTGAGGTGGCTCTGGACCTGTGGGAAGACTTCAGCCTCTGTAGAGAAGGGCAG	1125
Qy	261	ArgGluTrpValIleGlnLysIleHisGluSerGlnPheIleIleValValCysSerLys	280
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Qy	281	GlyMetLysTyrPheValAspLysLysAsnTyrLysHisLysGlyGlyGlyArgGlySer	300
Db	1186	GGTATGAAGTACTTTGTGGACAAGAAGAACTACAAACACAAAGGAGGTGGCCGAGGCTCG	1245
Qy	301	GlyLysGlyGluLeuPheLeuValAlaValSerAlaIleAlaGluLysLeuArgGlnAla	320
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Qy	381	ThrArgGlnGlySerArgArgAsnTyrPheArgSerLysSerGlyArgSerLeuTyrVal	400
Db	1486	ACGCGACAGGGCAGCAGAAGGAAGTACTTCCGGAGCAAGTCAGGCCGGTCCCTATACGTC	1545
Qy	401	AlaIleCysAsnMetHisGlnPheIleAspGluGluProAspTrpPheGluLysGlnPhe	420
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Qy	421	ValProPheHisProProProLeuArgTyrArgGluProValLeuGluLysPheAspSer	440
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Sequence
Comparison A

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Db      1726 GTAGAGGCGGCTGTTCTTGGGGCAACCGGACCAGCCGACTCCCAGCACGAGAGTCAGCAT 1785
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Qy      521 TyrAspSerSerValProSerSerGluLeuSerLeuProLeuMetGluGlyLeuSerThr 540
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Qy      541 AspGlnThrGluThrSerSerLeuThrGluSerValSerSerSerSerGlyLeuGlyGlu 560
Db      1966 GACCAGACAGAAACGTCTTCCCTGACGGAGAGCGTGTCTCTCTTCAGGCCTGGGTGAG 2025
Qy      561 GluGluProProAlaLeuProSerLysLeuLeuSerSerGlySerCysLysAlaAspLeu 580
Db      2026 GAGGAACCTCCTGCCCTTCTTCCAAGCTCCTCTCTTCTGGGTCATGCAAAGCAGATCTT 2085
Qy      581 GlyCysArgSerTyrThrAspGluLeuHisAlaValAlaProLeu 595
Db      2086 GGTGCGCGAGCTACACTGATGAAGCTCCACGCGGTGCCCCCTTTG 2130

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RESULT 8
AX364573

LOCUS AX364573 2383 bp DNA linear PAT 15-FEB-2002

DEFINITION Sequence 4 from Patent WO0208259.

ACCESSION AX364573

VERSION AX364573.1 GI:18696533

KEYWORDS

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1

AUTHORS Presnell, S.R., Kuestner, R.E. and Gao, Z.

TITLE Human cytokine receptor

JOURNAL Patent: WO 0208259-A 4 31-JAN-2002;

ZymoGenetics, Inc. (US)

FEATURES

source

Location/Qualifiers

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CDS

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